

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/537,847
Source: PCP/10
Date Processed by STIC: 2/2/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

10/539,847

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
Numbering The numbering under each 5th amino acid is misaligned. **Do not use tab codes between numbers; use space characters,** instead.
- 4 Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
"bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
- 7 Skipped Sequences
(OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
Response Per 1.823 of Sequence Rules, the **only valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
→ Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
"bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
TIME: 11:46:47

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

```

4 <110> APPLICANT: LIAU, Gene
5 STEFANSSON, Steingrimur
6 SU, Joseph
8 <120> TITLE OF INVENTION: ENDOTHELIAL CELL SPECIFICALLY BINDING
9 PEPTIDES
11 <130> FILE REFERENCE: 4-32422A/GTI/USN
13 <140> CURRENT APPLICATION NUMBER: 10/537,847
C--> 14 <141> CURRENT FILING DATE: 2005-06-08
16 <150> PRIOR APPLICATION NUMBER: PCT/EP03/014407
17 <151> PRIOR FILING DATE: 2003-12-17
19 <150> PRIOR APPLICATION NUMBER: 60/434,258
20 <151> PRIOR FILING DATE: 2002-12-18
22 <160> NUMBER OF SEQ ID NOS: 43
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 9
28 <212> TYPE: PRT
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 34 <221> NAME/KEY: PEPTIDE
35 <222> LOCATION: (1)...(9)
36 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 38 <400> 1
39 Cys Pro Asp Leu His His His Met Cys
40 1 5
43 <210> SEQ ID NO: 2
44 <211> LENGTH: 9
45 <212> TYPE: PRT
46 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
49 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 51 <221> NAME/KEY: PEPTIDE
52 <222> LOCATION: (1)...(9)
53 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 55 <400> 2
56 Cys Leu Gly Gln His Ala Phe Thr Cys
57 1 5
60 <210> SEQ ID NO: 3
61 <211> LENGTH: 9
62 <212> TYPE: PRT
63 <213> ORGANISM: Artificial Sequence
65 <220> FEATURE:

```

ppn-6
Does Not Comply
Corrected Diskette Needed
(give source of genetic material)

insufficient explanation

(see item 11)

on Error

summary

sheet)

same error

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
TIME: 11:46:47

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

66 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 68 <221> NAME/KEY: PEPTIDE
69 <222> LOCATION: (1)...(9)
70 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 72 <400> 3
73 Cys Ser Ser Asn Thr Ala Pro His Cys
74 1 5
77 <210> SEQ ID NO: 4
78 <211> LENGTH: 9
79 <212> TYPE: PRT
80 <213> ORGANISM: Artificial Sequence
82 <220> FEATURE:
83 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 85 <221> NAME/KEY: PEPTIDE
86 <222> LOCATION: (1)...(9)
87 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 89 <400> 4
90 Cys His Val Leu Pro Asn Gly Asn Cys
91 1 5
94 <210> SEQ ID NO: 5
95 <211> LENGTH: 10
96 <212> TYPE: PRT
97 <213> ORGANISM: Artificial Sequence
99 <220> FEATURE:
100 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 102 <221> NAME/KEY: PEPTIDE
103 <222> LOCATION: (1)...(9) (10)? since 10 amino acids are in the sequence
104 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 106 <400> 5
107 Cys Lys Pro Gly Ile Tyr Pro Ser Leu Cys
108 1 5 10
111 <210> SEQ ID NO: 6
112 <211> LENGTH: 9
113 <212> TYPE: PRT
114 <213> ORGANISM: Artificial Sequence
116 <220> FEATURE:
117 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 119 <221> NAME/KEY: PEPTIDE
120 <222> LOCATION: (1)...(9)
121 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 123 <400> 6
124 Cys Gln Thr Ala Arg Thr Pro Ala Cys
125 1 5
128 <210> SEQ ID NO: 7
129 <211> LENGTH: 9
130 <212> TYPE: PRT
131 <213> ORGANISM: Artificial Sequence
133 <220> FEATURE:
134 <223> OTHER INFORMATION: Artificial peptide sequence

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
TIME: 11:46:47

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

W--> 136 <221> NAME/KEY: PEPTIDE
137 <222> LOCATION: (1)...(9)
138 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 140 <400> 7
141 Cys Asn Gln Ser Gln Pro Lys His Cys
142 1 5
145 <210> SEQ ID NO: 8
146 <211> LENGTH: 9
147 <212> TYPE: PRT
148 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 153 <221> NAME/KEY: PEPTIDE
154 <222> LOCATION: (1)...(9)
155 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 157 <400> 8
158 Cys Thr Pro Ser Lys Ile Ser Val Cys
159 1 5
162 <210> SEQ ID NO: 9
163 <211> LENGTH: 9
164 <212> TYPE: PRT
165 <213> ORGANISM: Artificial Sequence
167 <220> FEATURE:
168 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 170 <221> NAME/KEY: PEPTIDE
171 <222> LOCATION: (1)...(9)
172 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 174 <400> 9
175 Cys Val Ser Pro Gly Pro Arg Leu Cys
176 1 5
179 <210> SEQ ID NO: 10
180 <211> LENGTH: 9
181 <212> TYPE: PRT
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 187 <221> NAME/KEY: PEPTIDE
188 <222> LOCATION: (1)...(9)
189 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 191 <400> 10
192 Cys Tyr Ala Leu Ser Gly Val Pro Cys
193 1 5
196 <210> SEQ ID NO: 11
197 <211> LENGTH: 9
198 <212> TYPE: PRT
199 <213> ORGANISM: Artificial Sequence
201 <220> FEATURE:
202 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 204 <221> NAME/KEY: PEPTIDE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
TIME: 11:46:47

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

205 <222> LOCATION: (1)...(9)
206 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 208 <400> 11
209 Cys Lys His Pro Pro Gln Pro Phe Cys
210 1 5
213 <210> SEQ ID NO: 12
214 <211> LENGTH: 9
215 <212> TYPE: PRT
216 <213> ORGANISM: Artificial Sequence
218 <220> FEATURE:
219 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 221 <221> NAME/KEY: PEPTIDE
222 <222> LOCATION: (1)...(9)
223 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 225 <400> 12
226 Cys His Gln Ser Lys Pro Leu Leu Cys
227 1 5
230 <210> SEQ ID NO: 13
231 <211> LENGTH: 9
232 <212> TYPE: PRT
233 <213> ORGANISM: Artificial Sequence
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 238 <221> NAME/KEY: PEPTIDE
239 <222> LOCATION: (1)...(9)
240 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 242 <400> 13
243 Cys Pro Gly Pro Phe Ser Asn Trp Cys
244 1 5
247 <210> SEQ ID NO: 14
248 <211> LENGTH: 9
249 <212> TYPE: PRT
250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 255 <221> NAME/KEY: PEPTIDE
256 <222> LOCATION: (1)...(9)
257 <223>. OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 259 <400> 14
260 Cys Pro His Lys Thr His Leu Pro Cys
261 1 5
264 <210> SEQ ID NO: 15
265 <211> LENGTH: 9
266 <212> TYPE: PRT
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 272 <221> NAME/KEY: PEPTIDE
273 <222> LOCATION: (1)...(9)

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
TIME: 11:46:47

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

274 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 276 <400> 15
277 Cys Val Phe Pro Leu Ser His Tyr Cys
278 1 5
281 <210> SEQ ID NO: 16
282 <211> LENGTH: 9
283 <212> TYPE: PRT
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 289 <221> NAME/KEY: PEPTIDE
290 <222> LOCATION: (1)...(9)
291 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 293 <400> 16
294 Cys Asn Met Ile Ala Pro Ser Ser Cys
295 1 5
298 <210> SEQ ID NO: 17
299 <211> LENGTH: 9
300 <212> TYPE: PRT
301 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
304 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 306 <221> NAME/KEY: PEPTIDE
307 <222> LOCATION: (1)...(9)
308 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 310 <400> 17
311 Cys Thr Leu Gly Met Gln Phe Gln Cys
312 1 5
315 <210> SEQ ID NO: 18
316 <211> LENGTH: 9
317 <212> TYPE: PRT
318 <213> ORGANISM: Artificial Sequence
320 <220> FEATURE:
321 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 323 <221> NAME/KEY: PEPTIDE
324 <222> LOCATION: (1)...(9)
325 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding
W--> 327 <400> 18
328 Cys Thr Asn Pro Thr Gly Met Leu Cys
329 1 5
332 <210> SEQ ID NO: 19
333 <211> LENGTH: 9
334 <212> TYPE: PRT
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: Artificial peptide sequence
W--> 340 <221> NAME/KEY: PEPTIDE
341 <222> LOCATION: (1)...(9)
342 <223> OTHER INFORMATION: Peptide Specific for Endothelial Cell Binding

Please correct
this type of error
in subsequent
sequences

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/02/2006
PATENT APPLICATION: US/10/537,847 TIME: 11:46:48

FYI
Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:43; Xaa Pos. 2,3,8.

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/537,847

DATE: 02/02/2006
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Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:34 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:38 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:51 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:68 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:72 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:85 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:89 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:102 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:106 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:119 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:123 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:136 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:140 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:153 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:157 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:170 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:174 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:187 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:191 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:204 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:208 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:221 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:225 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:238 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:242 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13
L:255 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:259 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:272 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:276 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:289 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:293 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:306 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:310 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:323 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:327 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:340 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:344 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:357 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:361 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:374 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:378 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:391 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:395 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:408 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:412 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:425 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

VERIFICATION SUMMARY
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TIME: 11:46:48

Input Set : A:\SEQLIST.TXT
Output Set: N:\CRF4\02022006\J537847.raw

L:429 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:442 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:446 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:459 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:463 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:476 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:480 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:27
L:493 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:497 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:510 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:514 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29
L:527 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:531 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:30
L:544 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
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L:561 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:565 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:32
L:578 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:582 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:33
L:595 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:599 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:34
L:612 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:616 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35
L:629 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:633 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:646 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:650 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:37
L:663 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:668 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:38
L:679 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:684 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:39
L:695 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:701 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:40
L:712 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:718 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:41
L:729 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:733 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:42
L:746 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:750 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:43
L:751 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0